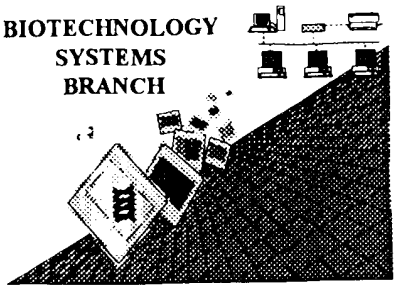
11-26-01

BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/581,528

Source: OIPE

Date Processed by STIC: 9/21/2001

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO **REDUCE** ERRORED SEQUENCE LISTINGS, **PLEASE** USE THE **CHECKER VERSION 3.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

http://www.uspto.gov/web/offices/pac/checker

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 09/581,528

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1	Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2	Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
3	Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
4	Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5	Variable Length	Sequence(s)____contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6	PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7	Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8	Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9	Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10	Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
11	Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12	PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13	Misuse of n	n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

AMC/MH – Biotechnology Systems Branch – 08/21/2001

OIPE

RAW SEQUENCE LISTING DATE: 09/21/2001
PATENT APPLICATION: **US/09/581,528** TIME: 18:15:30

Input Set : **A:\P19743.SL1.txt**
Output Set: **N:\CRF3\09212001\I581528.raw**

<110> APPLICANT: TAKEDA, Masatoshi
TAKEDA, Junji
<120> TITLE OF INVENTION: Gene Mutant Animals
<130> FILE REFERENCE: P19743
<140> CURRENT APPLICATION NUMBER: 09/581,528
<141> CURRENT FILING DATE: 1999-01-07
<150> PRIOR APPLICATION NUMBER: PCT/JP99/00015
<151> PRIOR FILING DATE: 1999-01-07
<160> NUMBER OF SEQ ID NOS: 17
<170> SOFTWARE: PatentIn version 3.0
<210> SEQ ID NO: 1
<211> LENGTH: 467
<212> TYPE: PRT
<213> ORGANISM: Human
<400> SEQUENCE: 1

Does Not Comply
Corrected Diskette Needed

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1               5                  10                  15
Ser Glu Asp Asn His Leu Ser Asn Thr Val Arg Ser Gln Asn Asp Asn
            20                  25                  30
Arg Glu Arg Gln Glu His Asn Asp Arg Arg Ser Leu Gly His Pro Glu
        35                  40                  45
Pro Leu Ser Asn Gly Arg Pro Gln Gly Asn Ser Arg Gln Val Val Glu
    50                  55                  60
Gln Asp Glu Glu Glu Asp Glu Glu Leu Thr Leu Lys Tyr Gly Ala Lys
65                  70                  75                  80
His Val Ile Met Leu Phe Val Pro Val Thr Leu Cys Met Val Val Val
                85                  90                  95
Val Ala Thr Ile Lys Ser Val Ser Phe Tyr Thr Arg Lys Asp Gly Gln
            100                 105                 110
Leu Ile Tyr Thr Pro Phe Thr Glu Asp Thr Glu Thr Val Gly Gln Arg
        115                 120                 125
Ala Leu His Ser Ile Leu Asn Ala Ala Ile Met Ile Ser Val Ile Val
    130                 135                 140
Val Met Thr Ile Leu Leu Val Val Leu Tyr Lys Tyr Arg Cys Tyr Lys
145                 150                 155                 160
Val Ile His Ala Trp Leu Ile Ile Ser Ser Leu Leu Leu Leu Phe Phe
                165                 170                 175
Phe Ser Phe Ile Tyr Leu Gly Glu Val Phe Lys Thr Tyr Asn Val Ala
            180                 185                 190
Val Asp Tyr Ile Thr Val Ala Leu Leu Ile Trp Asn Phe Gly Val Val
        195                 200                 205
Gly Met Ile Ser Ile His Trp Lys Gly Pro Leu Arg Leu Gln Gln Ala
    210                 215                 220
Tyr Leu Ile Met Ile Ser Ala Leu Met Ala Leu Val Phe Ile Lys Tyr
225                 230                 235                 240
Leu Pro Glu Trp Thr Ala Trp Leu Ile Leu Ala Val Ile Ser Val Tyr
                245                 250                 255
```

RAW SEQUENCE LISTING DATE: 09/21/2001
PATENT APPLICATION: **US/09/581,528** TIME: 18:15:30

Input Set : **A:\P19743.SL1.txt**
Output Set: **N:\CRF3\09212001\I581528.raw**

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75 Asp Leu Asp Ala Val Leu Cys Pro Lys Gly Pro Leu Arg Met Leu Val
76             260                 265                 270
78 Glu Thr Ala Gln Glu Arg Asn Glu Thr Leu Phe Pro Ala Leu Ile Tyr
79         275                 280                 285
81 Ser Ser Thr Met Val Trp Leu Val Asn Met Ala Glu Gly Asp Pro Glu
82     290                 295                 300
84 Ala Gln Arg Arg Val Ser Lys Asn Ser Lys Tyr Asn Ala Glu Ser Thr
85 305                 310                 315                 320
87 Glu Arg Glu Ser Gln Asp Thr Val Ala Glu Asn Asp Asp Gly Gly Phe
88                 325                 330                 335
90 Ser Glu Glu Trp Glu Ala Gln Arg Asp Ser His Leu Gly Pro His Arg
91             340                 345                 350
93 Ser Thr Pro Glu Ser Arg Ala Ala Val Gln Glu Leu Ser Ser Ser Ile
94         355                 360                 365
96 Leu Ala Gly Glu Asp Pro Glu Glu Arg Gly Val Lys Leu Gly Leu Gly
97     370                 375                 380
99 Asp Phe Ile Phe Tyr Ser Val Leu Val Gly Lys Ala Ser Ala Thr Ala
100 385                 390                 395                 400
102 Ser Gly Asp Trp Asn Thr Thr Ile Ala Cys Phe Val Ala Ile Leu Ile
103                 405                 410                 415
105 Gly Leu Cys Leu Thr Leu Leu Leu Leu Ala Ile Phe Lys Lys Ala Leu
106             420                 425                 430
108 Pro Ala Leu Pro Ile Ser Ile Thr Phe Gly Leu Val Phe Tyr Phe Ala
109         435                 440                 445
111 Thr Asp Tyr Leu Val Gln Pro Phe Met Asp Gln Leu Ala Phe His Gln
112     450                 455                 460
114 Phe Tyr Ile
115 465
117 <210> SEQ ID NO: 2
118 <211> LENGTH: 1404
119 <212> TYPE: DNA
120 <213> ORGANISM: Human
122 <400> SEQUENCE: 2
123 atgacagagt tacctgcacc gttgtcctac ttccagaatg cacagatgtc tgaggacaac     60
125 cacctgagca atactgtacg tagccagaat gacaatagag aacggcagga gcacaacgac    120
127 agacggagcc ttggccaccc tgagccatta tctaatggac gaccccaggg taactcccgg    180
129 caggtggtgg agcaagatga ggaagaagat gaggagctga cattgaaata tggcgccaag    240
131 catgtgatca tgctctttgt ccctgtgact ctctgcatgg tggtggtcgt ggctactatt    300
133 aagtcagtca gcttttatac ccggaaggat gggcagctaa tctatacccc attcacagaa    360
135 gataccgaga ctgtgggcca gagagccctg cactcaattc tgaatgctgc catcatgatc    420
137 agtgtcattg ttgtcatgac tatcctcctg gtggttctgt ataaatacag gtgctataag    480
139 gtcatccatg cctggcttat tatatcatct ctattgttgc tgttcttttt ttcattcatt    540
141 tacttggggg aagtgtttaa aacctataac gttgctgtgg actacattac tgttgcactc    600
143 ctgatctgga attttggtgt ggtgggaatg atttccattc actggaaagg tccacttcga    660
145 ctccagcagg catatctcat tatgattagt gccctcatgg ccctggtgtt tatcaagtac    720
147 ctccctgaat ggactgcgtg gctcatcttg gctgtgattt cagtatatga tttagtggct    780
149 gttttgtgtc cgaaaggtcc acttcgtatg ctggttgaaa cagctcagga gagaaatgaa    840
151 acgctttttc cagctctcat ttactcctca acaatggtgt ggttggtgaa tatggcagaa    900
153 ggagacccgg aagctcaaag gagagtatcc aaaaattcca agtataatgc agaaagcaca    960
```

RAW SEQUENCE LISTING DATE: 09/21/2001
PATENT APPLICATION: US/09/581,528 TIME: 18:15:30

Input Set : A:\P19743.SL1.txt
Output Set: N:\CRF3\09212001\I581528.raw

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155 gaaagggagt cacaagacac tgttgcagag aatgatgatg gcgggttcag tgaggaatgg   1020
157 gaagcccaga gggacagtca tctagggcct catcgctcta cacctgagtc acgagctgct   1080
159 gtccaggaac tttccagcag tatcctcgct ggtgaagacc cagaggaaag gggagtaaaa   1140
161 cttggattgg gagatttcat tttctacagt gttctggttg gtaaagcctc agcaacagcc   1200
163 agtggagact ggaacacaac catagcctgt ttcgtagcca tattaattgg tttgtgcctt   1260
165 acattattac tccttgccat tttcaagaaa gcattgccag ctcttccaat ctccatcacc   1320
167 tttgggcttg ttttctactt tgccacagat tatcttgtac agccttttat ggaccaatta   1380
169 gcattccatc aattttatat ctag                                         1404
172 <210> SEQ ID NO: 3
173 <211> LENGTH: 467
174 <212> TYPE: PRT
175 <213> ORGANISM: Mouse
177 <400> SEQUENCE: 3
179 Met Thr Glu Ile Pro Ala Pro Leu Ser Tyr Phe Gln Asn Ala Gln Met
180 1               5                  10                  15
182 Ser Glu Asp Ser His Ser Ser Ser Ala Ile Arg Ser Gln Asn Asp Ser
183             20                  25                  30
185 Glu Glu Arg Gln Gln Gln His Asp Arg Gln Arg Leu Asp Asn Pro Glu
186         35                  40                  45
188 Pro Ile Ser Asn Gly Arg Pro Gln Ser Asn Ser Arg Gln Val Val Glu
189     50                  55                  60
191 Gln Asp Glu Glu Glu Asp Glu Glu Leu Thr Leu Lys Tyr Gly Ala Lys
192 65                  70                  75                  80
194 His Val Ile Met Leu Phe Val Pro Val Thr Leu Cys Met Val Val Val
195                 85                  90                  95
197 Val Ala Thr Ile Lys Ser Val Ser Phe Tyr Thr Arg Lys Asp Gly Gln
198             100                 105                 110
200 Leu Ile Tyr Thr Pro Phe Thr Glu Asp Thr Glu Thr Val Gly Gln Arg
201         115                 120                 125
203 Ala Leu His Ser Ile Leu Asn Ala Ala Ile Met Ile Ser Val Ile Val
204     130                 135                 140
206 Ile Met Thr Ile Leu Leu Val Val Leu Tyr Lys Tyr Arg Cys Tyr Lys
207 145                 150                 155                 160
209 Val Ile His Ala Trp Leu Ile Ile Ser Ser Leu Leu Leu Leu Phe Phe
210                 165                 170                 175
212 Phe Ser Phe Ile Tyr Leu Gly Glu Val Phe Lys Thr Tyr Asn Val Ala
213             180                 185                 190
215 Val Asp Tyr Val Thr Val Ala Leu Leu Ile Trp Asn Phe Gly Val Val
216         195                 200                 205
218 Gly Met Ile Ala Ile His Trp Lys Gly Pro Leu Arg Leu Gln Gln Ala
219     210                 215                 220
221 Tyr Leu Ile Met Ile Ser Ala Leu Met Ala Leu Val Phe Ile Lys Tyr
222 225                 230                 235                 240
224 Leu Pro Glu Trp Thr Ala Trp Leu Ile Leu Ala Val Ile Ser Val Tyr
225                 245                 250                 255
227 Asp Leu Val Ala Val Leu Cys Pro Lys Gly Pro Leu Arg Met Leu Val
228             260                 265                 270
230 Glu Thr Ala Gln Glu Arg Asn Glu Thr Leu Phe Pro Ala Leu Ile Tyr
231         275                 280                 285
```

RAW SEQUENCE LISTING DATE: 09/21/2001
PATENT APPLICATION: US/09/581,528 TIME: 18:15:30

Input Set : A:\P19743.SL1.txt
Output Set: N:\CRF3\09212001\I581528.raw

```
233 Ser Ser Thr Met Val Trp Leu Val Asn Met Ala Glu Gly Asp Pro Glu
234     290                 295                 300
236 Ala Glu Arg Arg Val Pro Lys Asn Pro Lys Tyr Asn Thr Gln Arg Ala
237 305                 310                 315                 320
239 Glu Arg Glu Thr Gln Asp Ser Gly Ser Gly Asn Asp Asp Gly Gly Phe
240                 325                 330                 335
242 Ser Glu Glu Trp Glu Ala Gln Arg Asp Ser His Leu Gly Pro His Arg
243             340                 345                 350
245 Ser Thr Pro Glu Ser Arg Ala Ala Val Gln Glu Leu Ser Gly Ser Ile
246         355                 360                 365
248 Leu Thr Ser Glu Asp Pro Glu Glu Arg Gly Val Lys Leu Gly Leu Gly
249     370                 375                 380
251 Asp Phe Ile Phe Tyr Ser Val Leu Val Gly Lys Ala Ser Ala Thr Ala
252 385                 390                 395                 400
254 Ser Gly Asp Trp Asn Thr Thr Ile Ala Cys Phe Val Ala Ile Leu Ile
255                 405                 410                 415
257 Gly Leu Cys Leu Thr Leu Leu Leu Leu Ala Ile Phe Lys Lys Ala Leu
258             420                 425                 430
260 Pro Ala Leu Pro Ile Ser Ile Thr Phe Gly Leu Val Phe Tyr Phe Ala
261         435                 440                 445
263 Thr Asp Tyr Leu Val Gln Pro Phe Met Asp Gln Leu Ala Phe His Gln
264     450                 455                 460
266 Phe Tyr Ile
267 465
269 <210> SEQ ID NO: 4
270 <211> LENGTH: 1404
271 <212> TYPE: DNA
272 <213> ORGANISM: Mouse
274 <400> SEQUENCE: 4
275 atgacagaga tacctgcacc tttgtcctac ttccagaatg cccagatgtc tgaggacagc     60
277 cactccagca gcgccatccg gagccagaat gacagccaag aacggcagca gcagcatgac    120
279 aggcagagac ttgacaaccc tgagccaata tctaatgggc ggccccagag taactcaaga    180
281 caggtggtgg aacaagatga ggaggaagac gaagagctga cattgaaata tggagccaag    240
283 catgtcatca tgctctttgt ccccgtgacc ctctgcatgg tcgtcgtcgt ggccaccatc    300
285 aaatcagtca gcttctatac ccggaaggac ggtcagctaa tctacacccc attcacagaa    360
287 gacactgaga ctgtaggcca aagagccctg cactcgatcc tgaatgcggc catcatgatc    420
289 agtgtcattg tcattatgac catcctcctg gtggtcctgt ataaatacag gtgctacaag    480
291 gtcatccacg cctggcttat tatttcatct ctgttgttgc tgttcttttt ttcgttcatt    540
293 tacttagggg aagtatttaa gacctacaat gtcgccgtgg actacgttac agtagcactc    600
295 ctaatctgga attttggtgt ggtcgggatg attgccatcc actggaaagg ccccccttcga    660
297 ctgcagcagg cgtatctcat tatgatcagt gccctcatgg ccctggtatt tatcaagtac    720
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301 gttttatgtc ccaaaggccc acttcgtatg ctggttgaaa cagctcagga aagaaatgag    840
303 actctctttc cagctcttat ctattcctca acaatggtgt ggttggtgaa tatggctgaa    900
305 ggagacccag aagcccaaag gagggtaccc aagaacccca agtataacac acaaagagcg    960
307 gagagagaga cacaggacag tggttctggg aacgatgatg gtggcttcag tgaggagtgg   1020
309 gaggcccaaa gagacagtca cctggggcct catcgctcca ctcccgagtc aagagctgct   1080
311 gtccaggaac tttctgggag cattctaacg agtgaagacc cggaggaaag aggagtaaaa   1140
313 cttggactgg gagatttcat tttctacagt gttctggttg gtaaggcctc agcaaccgcc   1200
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RAW SEQUENCE LISTING DATE: 09/21/2001
PATENT APPLICATION: **US/09/581,528** TIME: 18:15:30

Input Set : **A:\P19743.SL1.txt**
Output Set: **N:\CRF3\09212001\I581528.raw**

315 agtggagact ggaacacaac catagcctgc tttgtagcca tactgatcgg cctgtgcctt 1260
317 acattactcc tgctcgccat tttcaagaaa gcgttgccag ccctccccat ctccatcacc 1320
319 ttcgggctcg tgttctactt cgccacggat taccttgtgc agcccttcat ggaccaactt 1380
321 gcattccatc agttttatat ctag 1404
324 <210> SEQ ID NO: 5
325 <211> LENGTH: 25
326 <212> TYPE: DNA
C--> 327 **<213> ORGANISM: Artificial**
329 <220> FEATURE:
330 <223> OTHER INFORMATION: Artificial Sequence
332 <400> SEQUENCE: 5
333 ggaattttgg tgtggtcggg atgat 25
336 <210> SEQ ID NO: 6
337 <211> LENGTH: 23
338 <212> TYPE: DNA
C--> 339 **<213> ORGANISM: Artificial**
341 <220> FEATURE:
342 <223> OTHER INFORMATION: Artificial Sequence
344 <400> SEQUENCE: 6
345 ggtccattcg gggaggtact tga 23
348 <210> SEQ ID NO: 7
349 <211> LENGTH: 36
350 <212> TYPE: DNA
C--> 351 **<213> ORGANISM: Artificial**
353 <220> FEATURE:
354 <223> OTHER INFORMATION: Artificial Sequence
356 <400> SEQUENCE: 7
357 tgtggtcggg atgatcgcca cccactggaa aggccc 36
360 <210> SEQ ID NO: 8
361 <211> LENGTH: 36
362 <212> TYPE: DNA
C--> 363 **<213> ORGANISM: Artificial**
365 <220> FEATURE:
366 <223> OTHER INFORMATION: Artificial Sequence
368 <400> SEQUENCE: 8
369 gggcctttcc agtgggtggc gatcatcccg accaca 36
372 <210> SEQ ID NO: 9
373 <211> LENGTH: 18
374 <212> TYPE: DNA
C--> 375 **<213> ORGANISM: Artificial**
377 <220> FEATURE:
378 <223> OTHER INFORMATION: Artificial Sequence
380 <400> SEQUENCE: 9
381 tctagacggc cgtctaga 18
384 <210> SEQ ID NO: 10
385 <211> LENGTH: 18
386 <212> TYPE: DNA
C--> 387 **<213> ORGANISM: Artificial**
389 <220> FEATURE:

insufficient response - give source of genetic material - see item 11 on Error Summary Sheet

VERIFICATION SUMMARY DATE: 09/21/2001
PATENT APPLICATION: **US/09/581,528** TIME: 18:15:31

Input Set : **A:\P19743.SL1.txt**
Output Set: **N:\CRF3\09212001\I581528.raw**

L:327 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
L:339 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
L:351 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
L:363 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
L:375 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
L:387 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
L:399 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
L:411 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12
L:423 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:13
L:435 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:447 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:459 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:471 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17